



PCT09

## RAW SEQUENCE LISTING

DATE: 08/28/2002

PATENT APPLICATION: US/09/830,328A

TIME: 12:46:59

Input Set : N:\Crf3\08262002\I830328.raw

Output Set: N:\CRF3\08282002\I830328A.raw

1 <110> APPLICANT: INCYTE PHARMACEUTICALS, INC.  
 2 TANG, Y. Tom  
 3 ARGENTINE, Charles C.  
 4 CORLEY, Neil C.  
 5 GORGONE, Gina A.  
 6 GUEGLER, Karl J.  
 7 BAUGHN, Mariah R.  
 8 <120> TITLE OF INVENTION: TRANSMEMBRANE 4 PROTEINS  
 9 <130> FILE REFERENCE: PF-0628 PCT  
 10 <140> CURRENT APPLICATION NUMBER: US/09/830,328A  
 11 <141> CURRENT FILING DATE: 2002-08-26  
 12 <150> PRIOR APPLICATION NUMBER: 09/183,027, unassigned  
 W--> 13 <151> PRIOR FILING DATE: 1998-10-29; 1998-10-29  
 14 <160> NUMBER OF SEQ ID NOS: 7  
 15 <170> SOFTWARE: PERL Program  
 17 <210> SEQ ID NO: 1  
 18 <211> LENGTH: 260  
 19 <212> TYPE: PRT  
 20 <213> ORGANISM: Homo sapiens  
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 22 <221> NAME/KEY: misc\_feature  
 23 <223> OTHER INFORMATION: Incyte ID No: 2651154CD1  
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 28 20 25 30  
 29 Ala Glu Cys Ile Phe Phe Val Ser Asp Gln His Ser Leu Tyr Pro  
 30 35 40 45  
 31 Leu Leu Glu Ala Thr Asp Asn Asp Asp Ile Tyr Gly Ala Ala Trp  
 32 50 55 60  
 33 Ile Gly Ile Phe Val Gly Ile Cys Leu Phe Cys Leu Ser Val Leu  
 34 65 70 75  
 35 Gly Ile Val Gly Ile Met Lys Ser Ser Arg Lys Ile Leu Leu Ala  
 36 80 85 90  
 37 Tyr Phe Ile Leu Met Phe Ile Val Tyr Ala Phe Glu Val Ala Ser  
 38 95 100 105  
 39 Cys Ile Thr Ala Ala Thr Gln Arg Asp Phe Phe Thr Pro Asn Leu  
 40 110 115 120  
 41 Phe Leu Lys Gln Met Leu Glu Arg Tyr Gln Asn Asn Ser Pro Pro  
 42 125 130 135  
 43 Asn Asn Asp Asp Gln Trp Lys Asn Asn Gly Val Thr Lys Thr Trp  
 44 140 145 150

Does Not Comply  
 Corrected Diskette Needed

pp 1,5

list prior application  
 number directly above  
 its filing date

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45   Asp Arg Leu Met Leu Gln Asp Asn Cys Cys Gly Val Asn Gly Pro
46                               155                      160          165
47   Ser Asp Trp Gln Lys Tyr Thr Ser Ala Phe Arg Thr Glu Asn Asn
48                               170                      175          180
49   Asp Ala Asp Tyr Pro Trp Pro Arg Gln Cys Cys Val Met Asn Asn
50                               185                      190          195
51   Leu Lys Glu Pro Leu Asn Leu Glu Ala Cys Lys Leu Gly Val Pro
52                               200                      205          210
53   Gly Phe Tyr His Asn Gln Gly Cys Tyr Glu Leu Ile Ser Gly Pro
54                               215                      220          225
55   Met Asn Arg His Ala Trp Gly Val Ala Trp Phe Gly Phe Ala Ile
56                               230                      235          240
57   Leu Cys Trp Thr Phe Trp Val Leu Leu Gly Thr Met Phe Tyr Trp
58                               245                      250          255
59   Ser Arg Ile Glu Tyr
60                               260
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65 <213> ORGANISM: Homo sapiens
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67 <221> NAME/KEY: misc_feature
68 <223> OTHER INFORMATION: Incyte ID No: 2674553CD1
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73               20              25              30
74   Val Ser Ala Trp Met Arg Asp Tyr Leu Asn Asn Val Leu Thr Leu
75               35              40              45
76   Thr Ala Glu Thr Arg Val Glu Glu Ala Val Ile Leu Thr Tyr Phe
77               50              55              60
78   Pro Val Val His Pro Val Met Ile Ala Val Cys Cys Phe Leu Ile
79               65              70              75
80   Ile Val Gly Met Leu Gly Tyr Cys Gly Thr Val Lys Arg Asn Leu
81               80              85              90
82   Leu Leu Leu Ala Trp Tyr Phe Gly Ser Leu Leu Val Ile Phe Cys
83               95             100             105
84   Val Glu Leu Ala Cys Gly Val Trp Thr Tyr Glu Gln Glu Leu Met
85              110             115             120
86   Val Pro Val Gln Trp Ser Asp Met Val Thr Leu Lys Ala Arg Met
87              125             130             135
88   Thr Asn Tyr Gly Leu Pro Arg Tyr Arg Trp Leu Thr His Ala Trp
89              140             145             150
90   Asn Phe Phe Gln Arg Glu Phe Lys Cys Cys Gly Val Val Tyr Phe
91              155             160             165
92   Thr Asp Trp Leu Glu Met Thr Glu Met Asp Trp Pro Pro Asp Ser
93              170             175             180
94   Cys Cys Val Arg Glu Phe Pro Gly Cys Ser Lys Gln Ala His Gln

```

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95          185          190          195
96  Glu Asp Leu Ser Asp Leu Tyr Gln Glu Gly Cys Gly Lys Lys Met
97          200          205          210
98  Tyr Ser Phe Leu Arg Gly Thr Lys Gln Leu Gln Val Leu Arg Phe
99          215          220          225
100  Leu Gly Ile Ser Ile Gly Val Thr Gln Ile Leu Ala Met Ile Leu
101          230          235          240
102  Thr Ile Thr Leu Leu Trp Ala Leu Tyr Tyr Asp Arg Arg Glu Pro
103          245          250          255
104  Gly Thr Asp Gln Met Met Ser Leu Lys Asn Asp Asn Ser Gln His
105          260          265          270
106  Leu Ser Cys Pro Ser Val Glu Leu Leu Lys Pro Ser Leu Ser Arg
107          275          280          285
108  Ile Phe Glu His Thr Ser Met Ala Asn Ser Phe Asn Thr His Phe
109          290          295          300
110  Glu Met Glu Glu Leu
111          305
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115 <212> TYPE: DNA
116 <213> ORGANISM: Homo sapiens
117 <220> FEATURE:
118 <221> NAME/KEY: misc_feature
119 <223> OTHER INFORMATION: Incyte ID No: 2651154CB1
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122  atggccaaag acaactcaac tgttcggtgc ttccagggcc tgctgatttt tggaaatgtg 120
123  attattgggt gttgcggcat tgccctgact gcggagtgc tcttctttgt atctgaccaa 180
124  cacagcctct acccactgct tgaagccacc gacaacgatg acatctatgg ggctgcctgg 240
125  atcggcata tttgtgggcat ctgcctcttc tgctgtctg ttctaggcat ttagggcatc 300
126  atgaagtcca gcaggaaaat tcttctggcg tatttcattc tgatgtttat agtatatgcc 360
127  tttgaagtgg catcttgtat cacagcagca acacaacgag actttttcac acccaacctc 420
128  ttctgaagc agatgctaga gaggtaccaa aacaacagcc ctccaaacaa tgatgaccag 480
129  tggaaaaaca atggagtcac caaaacctgg gacaggctca tgctccagga caattgctgt 540
130  ggcgtaaatg gtccatcaga ctggcaaaaa tacacatctg ccttccggac tgagaataat 600
131  gatgctgact atccctggcc tcgtcaatgc tgtgttatga acaatcttaa agaacctctc 660
132  aacctggagg cttgtaaact aggcgtgcct ggttttatc acaatcaggg ctgctatgaa 720
133  ctgatctctg gtccaatgaa ccgacacgcc tgggggggtg cctgggtttg atttgccatt 780
134  ctctgctgga ctttttgggt tctcctgggt accatgttct actggagcag aattgaatat 840
135  taagcataaa gtgttgccac catacctcct tccccgagtg actctggatt tgggtgctgga 900
136  accagctctc tccaatatt ccacgtttgt gccccacact aacgtgtgtg tottacattg 960
137  ccaagtcaga tggtagcgac ttcctttagg atctcaggct tctgcagttc tcatgactcc 1020
138  tacttttcat ctagtcttag cattctgcaa catttatata gactgttgaa aggagaattt 1080
139  gaaaaatgca taataacta ttccatccct gcttattttt aatttgggaa aataaataca 1140
140  ttcgaaggaa aaacaaaaaa aaggcgggcc cccgattatt gaggggtccc gagccgaac 1200
141  tcgtaaccat gtaaaacccg tccccgggg taaaattgta atccccccac aattccccaa 1260
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144          1331
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## RAW SEQUENCE LISTING

DATE: 08/28/2002

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TIME: 12:46:59

Input Set : N:\Crf3\08262002\I830328.raw

Output Set: N:\CRF3\08282002\I830328A.raw

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146 <211> LENGTH: 2768
147 <212> TYPE: DNA
148 <213> ORGANISM: Homo sapiens
149 <220> FEATURE:
150 <221> NAME/KEY: misc_feature
151 <223> OTHER INFORMATION: Incyte ID No: 2674553CB1
152 <400> SEQUENCE: 4
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155   gggacaagaa acaccacta ggacccaacc ccggcagcca gcggctcgag catgcgctga 180
156   gagtttgtgc agctggccct ggctgccgcc gctgcctcgt ccggactcgg agaggacttg 240
157   ggagggacag cggcgctggg aggtggctta gcagagactt tccagcaact gctgcccgag 300
158   actttttttt tttttttct ttttcccagg aggcggcgac ggcggcgggc gggggagagg 360
159   aagagaaaga agcgtctcca gctgaagcca atgcagccct ccggctctcc gcgaagaagt 420
160   tccctgcccc gatgagcccc cgccgtgcgt ccccgactat ccccgaggcg gcgtggggca 480
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162   atggggcttc tcccttacgg ggctcacaat ggccagagaa gattccgtga agtgtctgcg 600
163   ctgcctgctc tacgccctca atctgctctt ttggttaatg tccatcagtg tgttggcagt 660
164   ttctgcttgg atgagggact acctaaataa tgttctcact ttaactgcag aaacgagggt 720
165   agaggaagca gtcattttga cttactttcc tgtggttcat ccggtcatga ttgctgtttg 780
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168   cgtttggaca tatgaacagg aacttatggg tccagtacaa tggtcagata tggtcacttt 960
169   gaaagccagg atgacaaatt atggattacc tagatatcgg tggcttactc atgcttggaa 1020
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171   gacagagatg gactggcccc cagattcctg ctgtgttaga gaattcccag gatgttccaa 1140
172   acagcccac caggaagatc tcagtacact ttatcaagag gttgtggga agaaaatgta 1200
173   ttcttttttg agaggaacca aacaactgca ggtgctgagg tttctgggaa tctccatttg 1260
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175   tagaaggag cggggacag accaaatgat gtccttgaag aatgacaact ctgagcacct 1380
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178   gaagaaaacc acaaacttgt tttactggac ttgtgaattt ttgagtacat actatgtgtt 1560
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181   atgcccttaa aatgctgaag acagatgtca taccactgtg gtacgtgtg tatgactttt 1740
182   actgaacaca gttatgtttt gaggcagcat ggtttgatta gcatttccgc atccatgcaa 1800
183   acgagtcaca tatggtggga ctggagccat agtaaagggt gatttacttc taccaactag 1860
184   tatataaagt actaattaaa tgctaacata ggaagttaga aaatactaat aacttttatt 1920
185   actcagcgat ctattcttct gatgctaaat aaattatata tcagaaaact ttcaatattg 1980
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188   gtataattca gtcgatttca gttctgataa tgtaaagaat aaccattatg aaaaggaaaa 2160
189   tttgtcctgt atagcatcat tatttttagc ctttctgtt aataaagctt tactattctg 2220
190   tcttgggctt atattacaca tataactgtt atttaaatat ttaaccacta attttgaaaa 2280
191   ttaccagtgt gatacatagg aatcattatt cagaatgtag tctggtcttt aggaagtatt 2340
192   aataagaaaa tttgcacata acttagttga ttcagaaagg acttgtagtc tgtttttctc 2400
193   ccaaataaag actctttttg aactaaaca ctttttaaaa agcttatctt tgcttctcc 2460
194   aaacaagaag caatagtctc caagtcaata taaattctac agaaaatagt gttctttttc 2520

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## RAW SEQUENCE LISTING

DATE: 08/28/2002

PATENT APPLICATION: US/09/830,328A

TIME: 12:46:59

Input Set : N:\Crf3\08262002\I830328.raw

Output Set: N:\CRF3\08282002\I830328A.raw

195 tccagaaaaa tgcttgtag aatcattaaa acatgtgaca atttagagat tctttgtttt 2580  
 196 atttcactga ttaatatact gtggcaaatt acacagatta ttaaattttt ttacaagagt 2640  
 197 atagtatatatt tatttgaaat gggaaaagt cattttactg tattttgtgt attttgttta 2700  
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 199 aaaaaaaaaa 2768

201 &lt;210&gt; SEQ ID NO: 5

202 &lt;211&gt; LENGTH: 260

203 &lt;212&gt; TYPE: PRT

204 &lt;213&gt; ORGANISM: Bos taurus

205 &lt;300&gt; PUBLICATION INFORMATION:

206 &lt;308&gt; DATABASE ACCESSION NO: GenBank ID No: q443785

W--&gt; 207 &lt;300&gt; PUBLICATION INFORMATION: 5

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 212 Ala Glu Cys Ile Phe Phe Val Ser Asp Gln Asn Ser Leu Tyr Pro  
 213 35 40 45  
 214 Leu Leu Glu Ala Thr Asn Asn Asp Asp Ile Tyr Ala Ala Ala Trp  
 215 50 55 60  
 216 Ile Gly Met Ser Val Gly Ile Cys Leu Phe Cys Leu Ser Val Leu  
 217 65 70 75  
 218 Gly Ile Val Gly Ile Met Lys Ser Asn Arg Lys Ile Leu Leu Val  
 219 80 85 90  
 220 Tyr Phe Ile Leu Met Phe Ile Val Tyr Ala Phe Glu Val Ala Ser  
 221 95 100 105  
 222 Cys Ile Thr Ala Ala Thr Gln Arg Asp Phe Phe Thr Pro Asn Leu  
 223 110 115 120  
 224 Phe Leu Lys Gln Met Leu Glu Arg Tyr Gln Asn Asn Ser Pro Pro  
 225 125 130 135  
 226 Asn Asn Asp Asp Gln Trp Lys Asn Asn Gly Val Thr Lys Thr Trp  
 227 140 145 150  
 228 Asp Arg Leu Met Leu Gln Asp Asn Cys Cys Gly Val Asn Gly Pro  
 229 155 160 165  
 230 Ser Asp Trp Gln Lys Tyr Thr Ser Ala Phe Arg Thr Glu Asn Ser  
 231 170 175 180  
 232 Asp Ala Asp Tyr Pro Trp Pro Arg Gln Cys Cys Val Met Asn Ser  
 233 185 190 195  
 234 Leu Lys Glu Pro Leu Asn Leu Asp Ala Cys Lys Leu Gly Val Pro  
 235 200 205 210  
 236 Gly Tyr Tyr His Ser His Gly Cys Tyr Glu Leu Ile Ser Gly Pro  
 237 215 220 225  
 238 Met Asn Arg His Ala Trp Gly Val Ala Trp Phe Gly Phe Ala Ile  
 239 230 235 240  
 240 Leu Cys Trp Thr Phe Trp Val Leu Leu Gly Thr Met Phe Tyr Trp  
 241 245 250 255  
 242 Ser Arg Ile Asp Tyr  
 243 260  
 245 <210> SEQ ID NO: 6

> <309> insert this  
 and its  
 response whenever  
 <308> has a  
 response  
 same  
 error  
 in seqs 6-7

**VERIFICATION SUMMARY**

DATE: 08/28/2002

PATENT APPLICATION: US/09/830,328A

TIME: 12:47:00

Input Set : N:\Crf3\08262002\I830328.raw

Output Set: N:\CRF3\08282002\I830328A.raw

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L:207 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:5  
L:251 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:6  
L:293 M:256 W: Invalid Numeric Header Field, Identifier <309> Expected, SEQ:7